



#2 OIPE

RAW SEQUENCE LISTING
PATENT APPLICATION: US/10/010,229

DATE: 01/28/2002
TIME: 11:32:05

Input Set : N:\Crf3\RULE60\10010229.raw
Output Set: N:\CRF3\01282002\J010229.raw

1 <110> APPLICANT: Le, Junming
 2 Vilcek, Jan
 3 Daddona, Peter
 4 Ghrayeb, John
 5 Knight, David M.
 6 Siegel, Scott
 7 <120> TITLE OF INVENTION: Anti-TNF Antibodies and Peptides of
 8 Human Tumor Necrosis Factor
 9 <130> FILE REFERENCE: 0975.1005-013
 10 <140> CURRENT APPLICATION NUMBER: 10/010,229
 11 <141> CURRENT FILING DATE: 2001-12-07
 12 <150> PRIOR APPLICATION NUMBER: US/09/927,703
 13 <151> PRIOR FILING DATE: 2001-08-10
 15 <160> NUMBER OF SEQ ID NOS: 19
 16 <170> SOFTWARE: FastSEQ for Windows Version 4.0
 18 <210> SEQ ID NO: 1
 19 <211> LENGTH: 157
 20 <212> TYPE: PRT
 21 <213> ORGANISM: Homo sapiens
 22 <400> SEQUENCE: 1
 23 Val Arg Ser Ser Ser Arg Thr Pro Ser Asp Lys Pro Val Ala His Val
 24 1 5 10 15
 25 Val Ala Asn Pro Gln Ala Glu Gly Gln Leu Gln Trp Leu Asn Arg Arg
 26 20 25 30
 27 Ala Asn Ala Leu Leu Ala Asn Gly Val Glu Leu Arg Asp Asn Gln Leu
 28 35 40 45
 29 Val Val Pro Ser Glu Gly Leu Tyr Leu Ile Tyr Ser Gln Val Leu Phe
 30 50 55 60
 31 Lys Gly Gln Gly Cys Pro Ser Thr His Val Leu Leu Thr His Thr Ile
 32 65 70 75 80
 33 Ser Arg Ile Ala Val Ser Tyr Gln Thr Lys Val Asn Leu Leu Ser Ala
 34 85 90 95
 35 Ile Lys Ser Pro Cys Gln Arg Glu Thr Pro Glu Gly Ala Glu Ala Lys
 36 100 105 110
 37 Pro Trp Tyr Glu Pro Ile Tyr Leu Gly Gly Val Phe Gln Leu Glu Lys
 38 115 120 125
 39 Gly Asp Arg Leu Ser Ala Glu Ile Asn Arg Pro Asp Tyr Leu Asp Phe
 40 130 135 140
 41 Ala Glu Ser Gly Gln Val Tyr Phe Gly Ile Ile Ala Leu
 42 145 150 155
 44 <210> SEQ ID NO: 2
 45 <211> LENGTH: 321
 46 <212> TYPE: DNA

ENTERED

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47 <213> ORGANISM: Mus Balb/c
48 <220> FEATURE:
49 <221> NAME/KEY: CDS
50 <222> LOCATION: (1)...(321)
51 <400> SEQUENCE: 2
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      Asp Ile Leu Leu Thr Gln Ser Pro Ala Ile Leu Ser Val Ser Pro Gly
      1           5           10          15
      gaa aga gtc agt ttc tcc tgc agg gcc agt cag ttc gtt ggc tca agc  96
      Glu Arg Val Ser Phe Ser Cys Arg Ala Ser Gln Phe Val Gly Ser Ser
      20          25          30
      atc cac tgg tat cag caa aga aca aat ggt tct cca agg ctt ctc ata 144
      Ile His Trp Tyr Gln Gln Arg Thr Asn Gly Ser Pro Arg Leu Leu Ile
      35          40          45
      aag tat gct tct gag tct atg tct ggg atc cct tcc agg ttt agt ggc 192
      Lys Tyr Ala Ser Glu Ser Met Ser Gly Ile Pro Ser Arg Phe Ser Gly
      50          55          60
      agt gga tca ggg aca gat ttt act ctt agc atc aac act gtg gag tct 240
      Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn Thr Val Glu Ser
      65          70          75          80
      gaa gat att gca gat tat tac tgt caa caa agt cat agc tgg cca ttc 288
      Glu Asp Ile Ala Asp Tyr Tyr Cys Gln Gln Ser His Ser Trp Pro Phe
      85          90          95
      acg ttc ggc tcg ggg aca aat ttg gaa gta aaa 321
      Thr Phe Gly Ser Gly Thr Asn Leu Glu Val Lys
      100         105
74 <210> SEQ ID NO: 3
75 <211> LENGTH: 107
76 <212> TYPE: PRT
77 <213> ORGANISM: Mus Balb/c
78 <400> SEQUENCE: 3
      Asp Ile Leu Leu Thr Gln Ser Pro Ala Ile Leu Ser Val Ser Pro Gly
      1           5           10          15
      Glu Arg Val Ser Phe Ser Cys Arg Ala Ser Gln Phe Val Gly Ser Ser
      20          25          30
      Ile His Trp Tyr Gln Gln Arg Thr Asn Gly Ser Pro Arg Leu Leu Ile
      35          40          45
      Lys Tyr Ala Ser Glu Ser Met Ser Gly Ile Pro Ser Arg Phe Ser Gly
      50          55          60
      Ser Gly Ser Gly Thr Asp Phe Thr Leu Ser Ile Asn Thr Val Glu Ser
      65          70          75          80
      Glu Asp Ile Ala Asp Tyr Tyr Cys Gln Gln Ser His Ser Trp Pro Phe
      85          90          95
      Thr Phe Gly Ser Gly Thr Asn Leu Glu Val Lys
      100         105
94 <210> SEQ ID NO: 4
95 <211> LENGTH: 357
96 <212> TYPE: DNA
97 <213> ORGANISM: Mus Balb/c

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98 <220> FEATURE:
 99 <221> NAME/KEY: CDS
 100 <222> LOCATION: (1)...(357)
 101 <400> SEQUENCE: 4
 102 gaa gtg aag ctt gag gag tct gga gga ggc ttg gtg caa cct gga gga 48
 103 Glu Val Lys Leu Glu Glu Ser Gly Gly Leu Val Gln Pro Gly Gly
 104 1 5 10 15
 105 tcc atg aaa ctc tcc tgt gtt gcc tct gga ttc att ttc agt aac cac 96
 106 Ser Met Lys Leu Ser Cys Val Ala Ser Gly Phe Ile Phe Ser Asn His
 107 20 25 30
 108 tgg atg aac tgg gtc cgc cag tct cca gag aag ggg ctt gag tgg gtt 144
 109 Trp Met Asn Trp Val Arg Gln Ser Pro Glu Lys Gly Leu Glu Trp Val
 110 35 40 45
 111 gct gaa att aga tca aaa tct att aat tct gca aca cat tat gcg gag 192
 112 Ala Glu Ile Arg Ser Lys Ser Ile Asn Ser Ala Thr His Tyr Ala Glu
 113 50 55 60
 114 tct gtg aaa ggg agg ttc acc atc tca aga gat gat tcc aaa agt gct 240
 115 Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Ser Ala
 116 65 70 75 80
 117 gtc tac ctg caa atg acc gac tta aga act gaa gac act ggc gtt tat 288
 118 Val Tyr Leu Gln Met Thr Asp Leu Arg Thr Glu Asp Thr Gly Val Tyr
 119 85 90 95
 120 tac tgt tcc agg aat tac tac ggt agt acc tac gac tac tgg ggc caa 336
 121 Tyr Cys Ser Arg Asn Tyr Tyr Gly Ser Thr Tyr Asp Tyr Trp Gly Gln
 122 100 105 110
 123 ggc acc act ctc aca gtc tcc 357
 124 Gly Thr Thr Leu Thr Val Ser
 125 115
 127 <210> SEQ ID NO: 5
 128 <211> LENGTH: 119
 129 <212> TYPE: PRT
 130 <213> ORGANISM: Mus Balb/c
 131 <400> SEQUENCE: 5
 132 Glu Val Lys Leu Glu Glu Ser Gly Gly Leu Val Gln Pro Gly Gly
 133 1 5 10 15
 134 Ser Met Lys Leu Ser Cys Val Ala Ser Gly Phe Ile Phe Ser Asn His
 135 20 25 30
 136 Trp Met Asn Trp Val Arg Gln Ser Pro Glu Lys Gly Leu Glu Trp Val
 137 35 40 45
 138 Ala Glu Ile Arg Ser Lys Ser Ile Asn Ser Ala Thr His Tyr Ala Glu
 139 50 55 60
 140 Ser Val Lys Gly Arg Phe Thr Ile Ser Arg Asp Asp Ser Lys Ser Ala
 141 65 70 75 80
 142 Val Tyr Leu Gln Met Thr Asp Leu Arg Thr Glu Asp Thr Gly Val Tyr
 143 85 90 95
 144 Tyr Cys Ser Arg Asn Tyr Tyr Gly Ser Thr Tyr Asp Tyr Trp Gly Gln
 145 100 105 110
 146 Gly Thr Thr Leu Thr Val Ser
 147 115

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149 <210> SEQ ID NO: 6
150 <211> LENGTH: 8
151 <212> TYPE: PRT
152 <213> ORGANISM: Homo sapiens
153 <400> SEQUENCE: 6
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      1           5
157 <210> SEQ ID NO: 7
158 <211> LENGTH: 7
159 <212> TYPE: PRT
160 <213> ORGANISM: Homo sapiens
161 <400> SEQUENCE: 7
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      1           5
165 <210> SEQ ID NO: 8
166 <211> LENGTH: 20
167 <212> TYPE: DNA
168 <213> ORGANISM: Artificial Sequence
169 <220> FEATURE:
170 <223> OTHER INFORMATION: PCR oligonucleotides
171 <400> SEQUENCE: 8
      cctggataacc tgtgaaaaga
172                                         20
174 <210> SEQ ID NO: 9
175 <211> LENGTH: 27
176 <212> TYPE: DNA
177 <213> ORGANISM: Artificial Sequence
178 <220> FEATURE:
179 <223> OTHER INFORMATION: PCR oligonucleotides
180 <400> SEQUENCE: 9
      cctggtacct tagtcaccgt ctccctca
181                                         27
183 <210> SEQ ID NO: 10
184 <211> LENGTH: 27
185 <212> TYPE: DNA
186 <213> ORGANISM: Artificial Sequence
187 <220> FEATURE:
188 <223> OTHER INFORMATION: PCR oligonucleotides
189 <400> SEQUENCE: 10
      aatagatatac tccttcaaca cctgcaa
190                                         27
192 <210> SEQ ID NO: 11
193 <211> LENGTH: 21
194 <212> TYPE: DNA
195 <213> ORGANISM: Artificial Sequence
196 <220> FEATURE:
197 <223> OTHER INFORMATION: PCR oligonucleotides
198 <400> SEQUENCE: 11
      atcggggacaa agttggaaat a
199                                         21
201 <210> SEQ ID NO: 12
202 <211> LENGTH: 16
203 <212> TYPE: DNA

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Input Set : N:\Crf3\RULE60\10010229.raw
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204 <213> ORGANISM: Artificial Sequence
205 <220> FEATURE:
206 <223> OTHER INFORMATION: PCR oligonucleotides
207 <400> SEQUENCE: 12
208 ggcggtctgg taccgg 16
210 <210> SEQ ID NO: 13
211 <211> LENGTH: 19
212 <212> TYPE: DNA
213 <213> ORGANISM: Artificial Sequence
214 <220> FEATURE:
215 <223> OTHER INFORMATION: PCR oligonucleotides
216 <400> SEQUENCE: 13
217 gtcaacaaca tagtcatca 19
219 <210> SEQ ID NO: 14
220 <211> LENGTH: 23
221 <212> TYPE: DNA
222 <213> ORGANISM: Artificial Sequence
223 <220> FEATURE:
224 <223> OTHER INFORMATION: PCR oligonucleotides
225 <400> SEQUENCE: 14
226 cacaggtgtg tccccaaaggaaa 23
228 <210> SEQ ID NO: 15
229 <211> LENGTH: 18
230 <212> TYPE: DNA
231 <213> ORGANISM: Artificial Sequence
232 <220> FEATURE:
233 <223> OTHER INFORMATION: PCR oligonucleotides
234 <400> SEQUENCE: 15
235 aatctggggtaggcacaa 18
237 <210> SEQ ID NO: 16
238 <211> LENGTH: 17
239 <212> TYPE: DNA
240 <213> ORGANISM: Artificial Sequence
241 <220> FEATURE:
242 <223> OTHER INFORMATION: PCR oligonucleotides
243 <400> SEQUENCE: 16
244 agtgtgtgtcccaagg 17
246 <210> SEQ ID NO: 17
247 <211> LENGTH: 24
248 <212> TYPE: DNA
249 <213> ORGANISM: Artificial Sequence
250 <220> FEATURE:
251 <223> OTHER INFORMATION: PCR oligonucleotides
252 <400> SEQUENCE: 17
253 cacagctgcc cgccccagggtgcata 24
255 <210> SEQ ID NO: 18
256 <211> LENGTH: 17
257 <212> TYPE: DNA
258 <213> ORGANISM: Artificial Sequence

VERIFICATION SUMMARY

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Input Set : N:\Crf3\RULE60\10010229.raw
Output Set: N:\CRF3\01282002\J010229.raw